

Fg. I

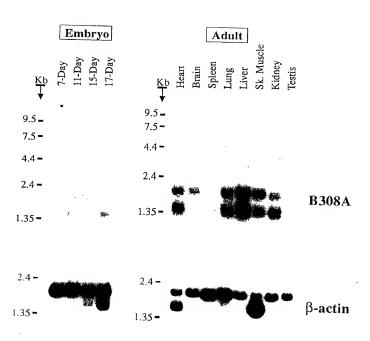
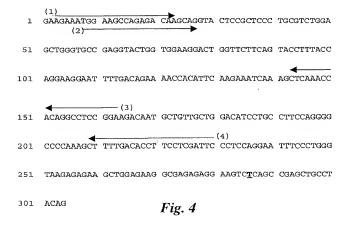


Fig. 2

TTGCCCTCAA CAAAGATGGT CTTTATGGTA CAGGTTCCCT AGCAGTCTGG ATTCCGGTTG TAGTTTTAGT TATTCTTTTT TTTTTTTTT TAAACGGTAC 101 GTGGTCGCAG ACGAAGAAAT GGAAGCCAGA GACAAGCAGG TACTCCGCTC CCTGCGTCTG GAGCTGGGTG CCGAGGTACT GGTGGAAGGA CTGGTTCTTC 151 201 AGTACCTTTA CCAGGAAGGA ATTTTGACAG AAAACCACAT TCAAGAAATC 251 AAAGCTCAAA CCACAGGCCT CCGGAAGACA ATGCTGTTGC TGGACATCCT GCCTTCCAGG GGCCCCAAAG CTTTTGACAÇ CTTCCTCGAT TCCCTCCAGG 301 351 AATTTCCCTG GGTAAGAGAG AAGCTGGAGA AGGCGAGAGA GGAAGTCTCA 401 GCCGAGCTGC CTACAGGTGA CTGGATGGCC GGAATCCCCT CACACATCCT CAGCAGCTCG CCATCAGACC AGCAGATTAA CCAGCTGGCT CAGAGGCTAG 451 GCCCGGAGTG GGAGCCCGTG GTCCTGTCTC TGGGACTGTC CCAGACCGAC 501 551 ATCTACCGCT GCAAGGCCAA CCATCCCCAC AACGTGCATT CGCAGGTGGT 601 GGAGGCCTTT GTCCGCTGGC GCCAGCGTTT TGGGAAGCAG GCCACCTTCC 651 TAAGCTTACA CAAGGGCCTC CAGGCAATGG AGGCTGATCC CTCCCTGCTC CAGCACATGC tGGAGTGACC TGACCCCCC CCGCGCCCCC CCCCCACTTG 701 751 CTGTGGGGGT GGTGGGGCGT GGGTTCCCAA GTCACACTGG CTGAACCGGA 801 CTTTTCTCAG CAGGTGGCTT TGTTCTGGGC TTTTCAGTGA TCTGTTTACG 851 GAAAGAGATC GTCCACCACT CACTCAACCA TCGATTGGCT TTAATTGCTT 901 GAAGACTGCG CTGTTGTAAC TATGGTTTGG AACTTTGTGG CTGGCCTTTA ACAGGAGGCC AGAAAAAACA CAACACCCAC CCTACCCAAC CCCCCAAAAA 951 1001 ATCATGCTAC AGCATCGAAT GCAGGTGTCC TGCATACAAG GCAGCTACAC 1051 TTGTGTTGCC TGGAGACTGG ATTGTGCATT TAGCTCTTCA TAATGGTGAT 1101 GATAATAAA AAGCAAATTG TGATATAGAA TGTGCCTCTT TCAATGAGAG 1201 CACACCAATC TTCTGTTGCA TAGACGGAGG GTGTAAAAAT ATGGGAGTGG 1251 AGCAAGATTG ATAGCAGTCA TGTGACGACG GAGATAAATA ACTCAGGCAG 1301 GATGTATAGA TTAAGCATGA GACACCGAAG CTCCCTGCAG AGGCCAGGGA 1351 GAGAACGGAA GACCTTCATC TTAACAAATT GTATGAGGAG TCTCTGTCCA 1401 TTTGTTAAAG GCATTGGATC AGAGACAAGA GGGCTCAGTG TTTCTCTTGA 1451 GGCCTGAATG GCTGAAGGCG GTGAGTTCCC GAGGGGCGTC ATGGGTTGTC 1501 CAGCCTTTCA TTAACTGCAC ATAGTGTTAG CCAGACAGGT GTACGTGTTT 1551 GTCATCCCAT CTAAGAGACT GAAGCAGGAG GATCACCTGT ACATGACTGC 1601 TTCTTTCAAC ATTTTAAAAT GTGTAACTTC TATTAAATTC TCTCAGTGCA 1651 AAAAAAAAA AAAAAAA Fig. 3A

MEARDKQVLRSLRLELGAEVLVEGLVLQYLYQEGILTENHIQEIKAQTTG LRKTMLLLDILPSRGPKAPDTFLDSLQBFPWVEKLEKAREBVSAELPTG DMMAGIPSHILSSSPSDQQINQLAQRLGPEWEPVVLSLGLSQTDIYRCKA NHPHNVHSOVVEAFVRWRORFGKQATFLSLHKGLQAMEADPSLLOHMLE"

## Fig. 3B



- 1 ggaaatggag gctagagaca agcaagtgct tcgctccctt cgcctggagt
- 51 tgggtgcaga ggtactggtg gaggggctag tcctccagta tctttatcag
- 101 gaaggggtct tgacagaaag ccacgttcaa gaaattaaag ctcaagccac
- 151 aggcctccgg

Fig. 5

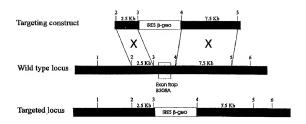


Fig.6

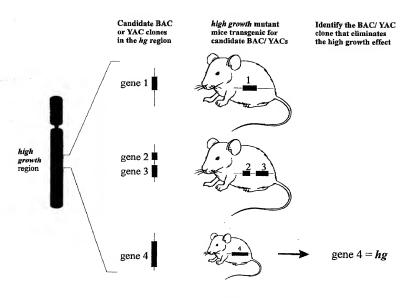


Fig. 7

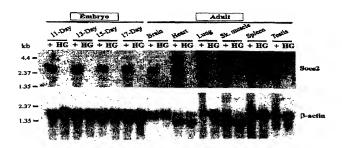


Fig. 8

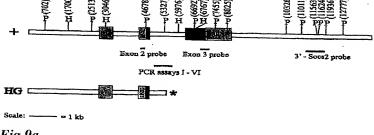


Fig 9a

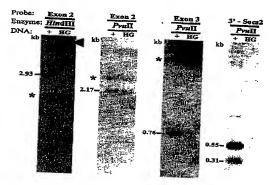


Fig 9b

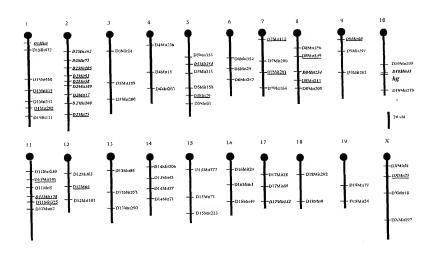
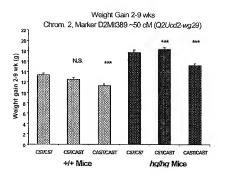
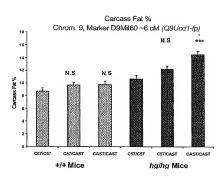
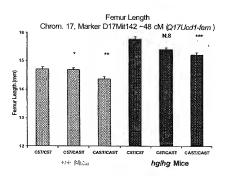


Fig. 10







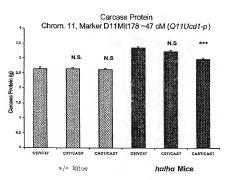


Fig. 11

## A: hg/hg mice

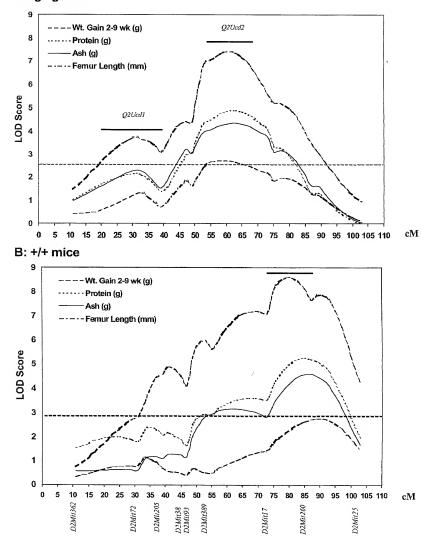


Fig. 12

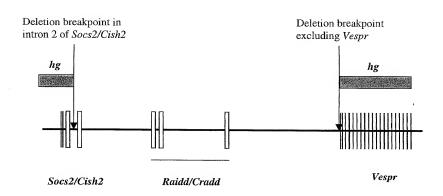


Fig. 13

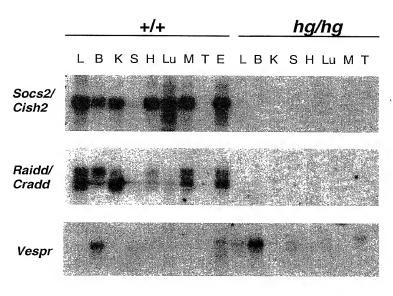


Fig. 14